

Figure 1

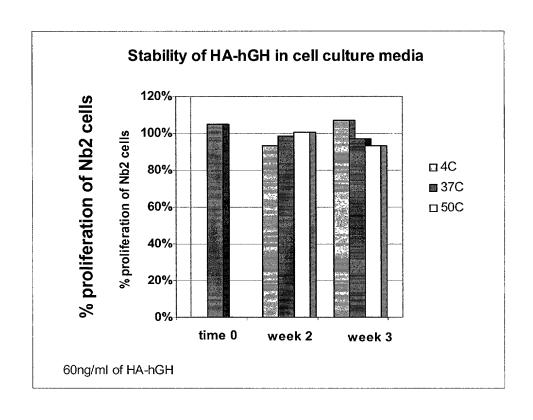


Figure 2

3/18

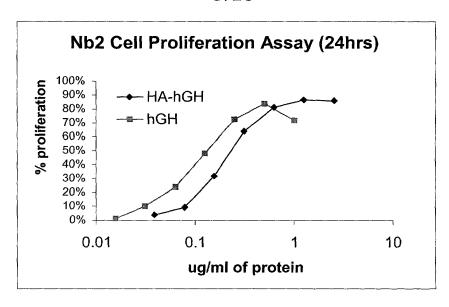


Figure 3A

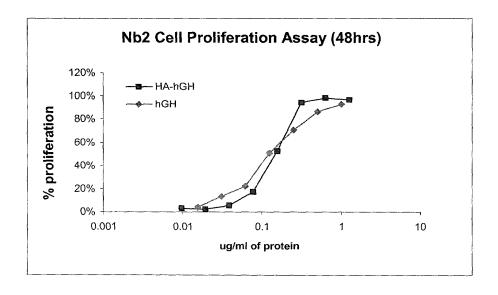


Figure 3B

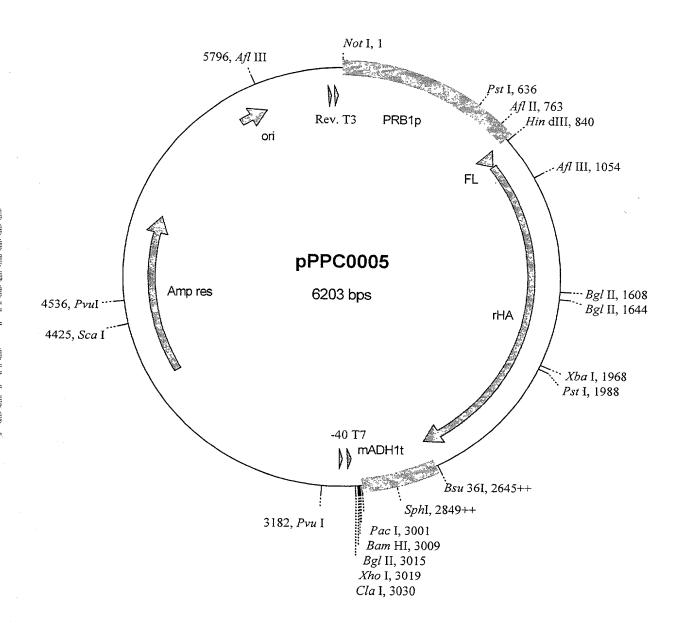


Figure 4

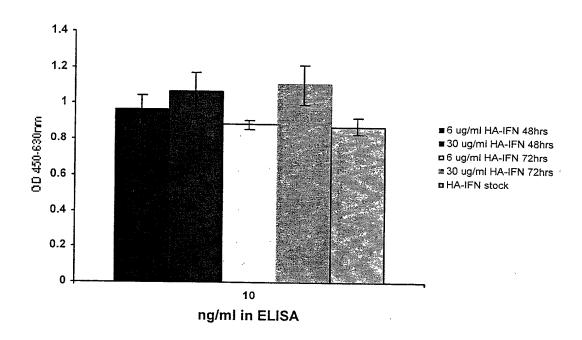
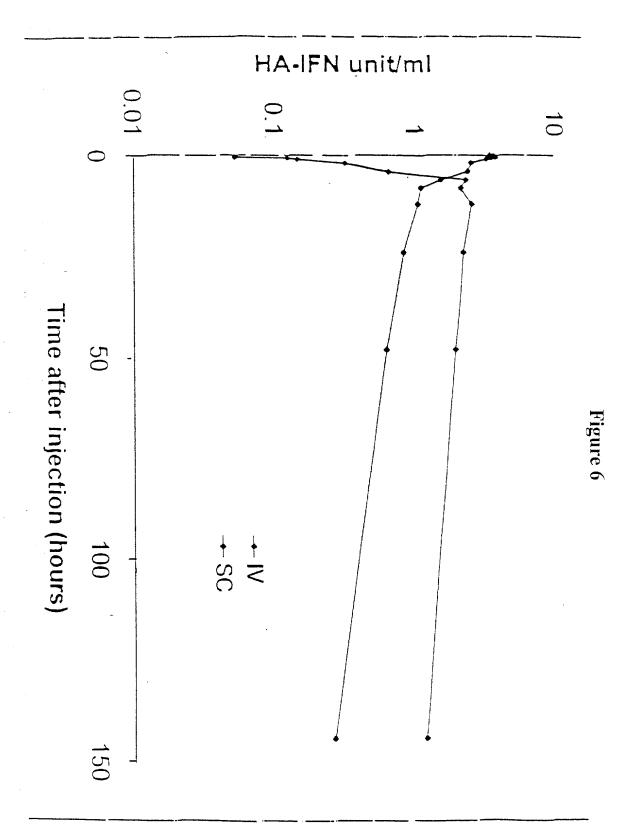


Figure 5



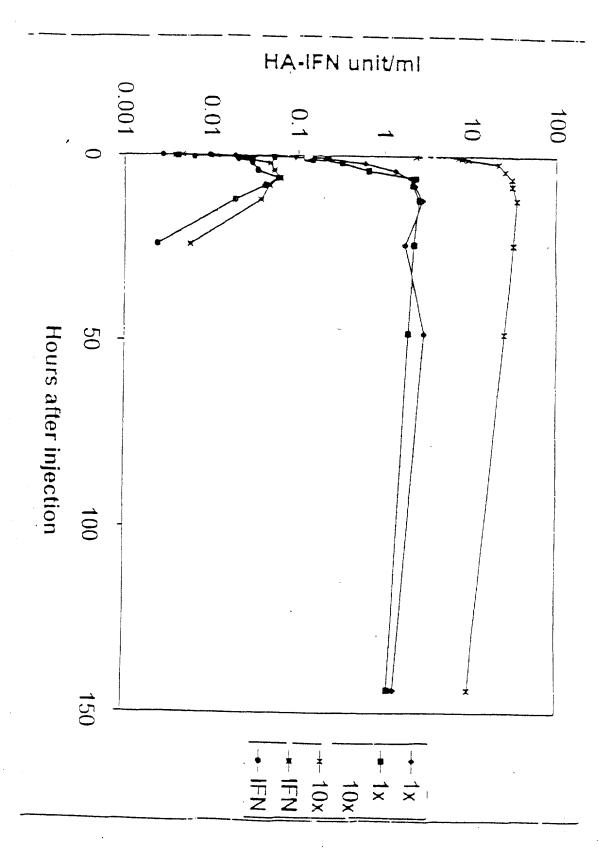


Figure 7

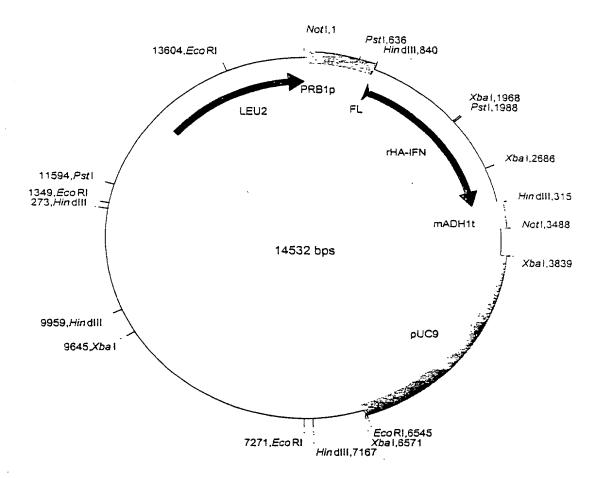


Figure 8. The HA-IFN α expression cassette in pSAC35. The expression cassette comprises

PRB1 promoter, from S. cerevisiae.

Fusion leader, first 19 amino acids of the HA leader followed by the last 6 amino acids of the MF α -1 leader.

HA-IFNα coding sequence with a double stop codon (TAATAA)

ADH1 terminator, from S. cerevisiae. Modified to remove all the coding sequence normaly present in the Hind III/BamHI fragment generally used.

Figure 8

Localisation of 'Loops' based on the HA Crystal Structure which could be used for Mutation/Insertion

1	DAHKSEVAHR HHHHH	FKDLGEENFK HHH HHH	ALVLIAFAQY ННННННННН	LQQCPFEDHV HHHHH	KLVNEVTEFA ННИНННННН
	I II III				
51	KTCVADESAE	NCDKSLHTLF	GDKLC TVATL	RETYGEMADC	
	ннннн	ннннн	ниннн	нннн	
101	CELOHKDDNP	NI.PRI.VR PEV	DVMCTAFHDN	EETFLKKYLY	FTADDUDVEV
	НННН			нинининн	
					
IV 151 APELLFFAKR YKAAFTECCO AADKAACLLP KLDELRDEGK ASSAKORLKC					
T 2 T				HHHEHHHHHH	
		***************************************	**********		
201	a de ouedena	T1111111111111111111111111111111111111			v
20I	ASLQKFGERA			VSKLVTDLTK HHHHHHHHHH	
	111111111111111111111111111111111111111			nnnnnnnn	nnnnn nn
	VI		VII		
251	LE CADDRADL	AKYIC ENODS	<u> ISSKLKECCE</u>	KPLLEKSHCI	AEVENDEMPA
	нинининни	нннн	ннннн	нннннн	Н
301	DLPSLAADFV	ESKDVCKNYA	EAKDVFLGMF	LYEYARRHPD	YSVVLLLRLA
	нннн	нннннн	нннннн	ннннн	ннннннн
VIII					
351	KTYETTLEKC		AKVFDEFKPL	VEEPONLIKO	NCELFEOLGE
	нннннннн	HH	н ннннн	нинининин	нниннин
401	YKFQNALLVR	VTKK1/DOMET	DTT TETTED MT	CVTCCVCCVI	IX
101				HHH	
451	X XI DYLSVVLNQL CVLHEKTPVS DRVTKCCTES LVNRRPPCFSA LEVDETYVE				
42T				HHHHHHHH	
501	EFNAETFTFH	ADICTLSEKE	RQIKKQTALV	ELVKHKPKAT	KEQLKAVMDD
		нни ннн	нннименнн	ННН	ннннннн
XII					
551			EGKKLVAASQ		
	ННННННН		нининнин	HH	
	Loop		Loop		
	I Val54-Asn61		VII	Glu280-His288	
	II Thr76-Asp89 III Ala92-Glu100		VIII IX	Ala362-Glu368 Lys439-Pro447	
	IV Gln170-Ala176		X	Lys439-Pr0447 Val462-Lys475	
		47-Glu252	XI	Thr478-Pro4	186
	VI Glu266-Glu277		XII	Lys560-Thr566	

Figure 9

Examples of Modifications to Loop IV

a. Randomisation of Loop IV.

ΙV

IV

X represents the mutation of the natural amino acid to any other amino acid. One, more or all of the amino acids can be changed in this manner. This figure indicates all the residues have been changed.

b. Insertion (or replacement) of Randomised sequence into Loop IV.

 $(X)_n$



IV

The insertion can be at any point on the loop and the length a length where n would typically be 6, 8, 12, 20 or 25.

Figure 10

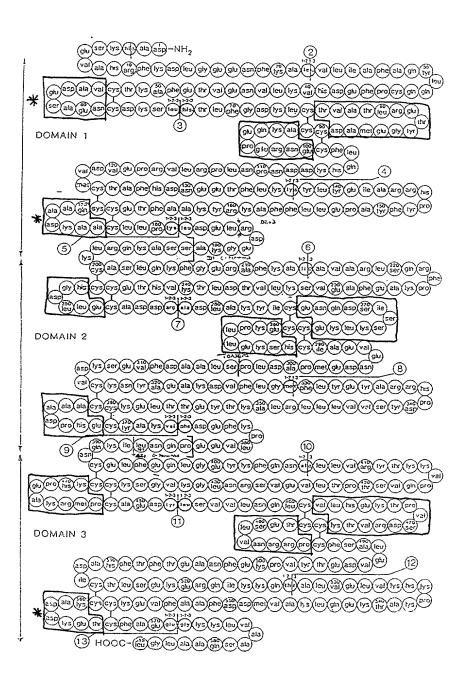
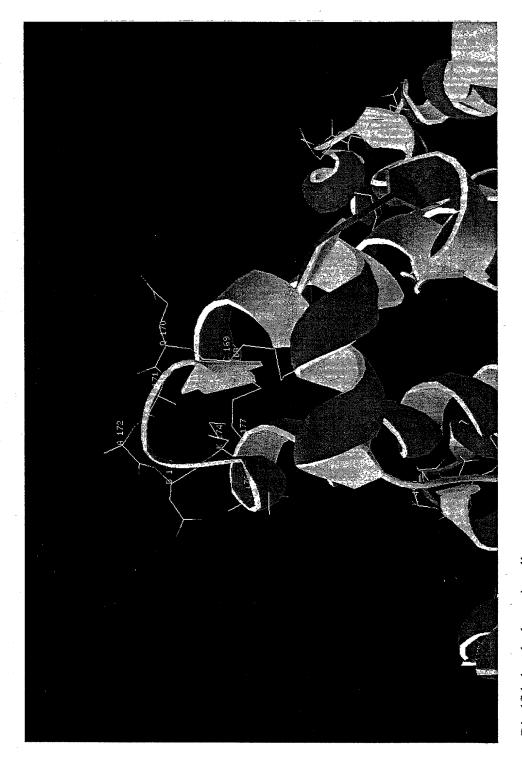


Figure 11



Disulfide bonds shown in yellow

Figure 12: Loop IV Gln170-Ala176

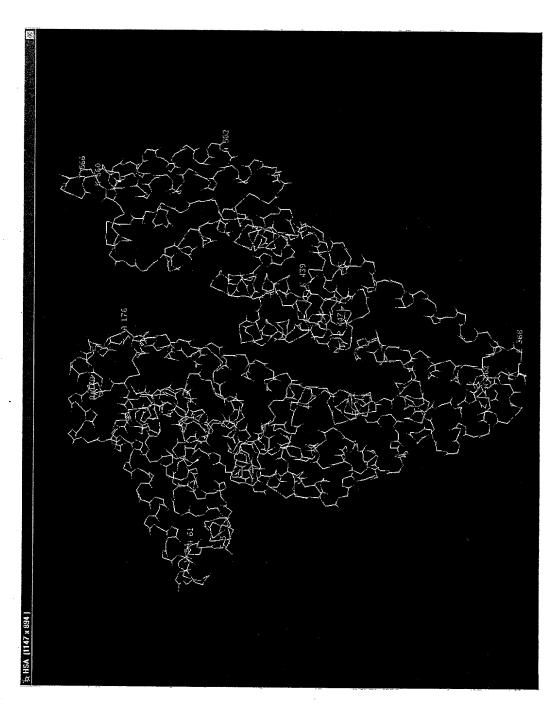
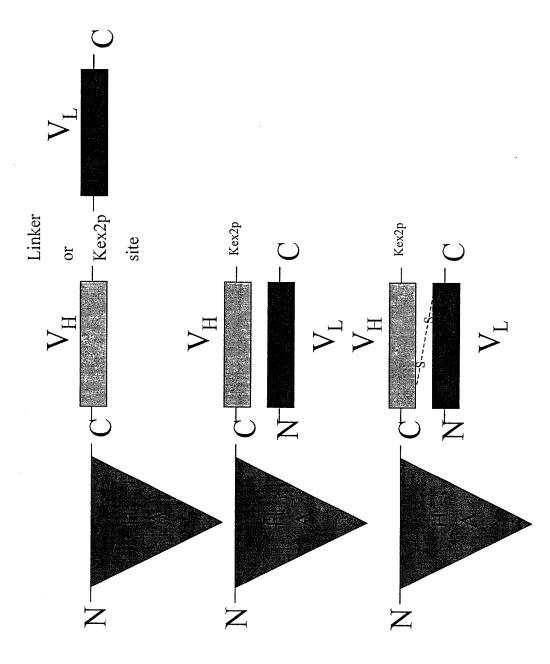


Figure 13: Tertiary Structure of HA



The first and the state of the

Figure 14: Schematic Diagram of Possible ScFv Fusions (Example is of a C-terminal fusion to HA)

GAA 180 E 60 240 80 GAA ATT GCC AGA AGA CAT CCT TAC TTT TAT GCC CCG GAA CTC CTT TTC TTT GCT AAA AGG 480 E I A R R H P Y F Y A P E L L F F A K R 160 CCT GAG AGA AAT GAA 300 P E R N E 100 TGC TTC TTG CAA CAC AAA GAT GAC AAC CCA AAC CTC CCC CGA TTG GTG AGA CCA GAG GTT 360 C F L Q H K D D N P N L P R L V R P E V 120 TTT TTG AAA AAA TAC TTA TAT 420 F L K K Y L Y 140 TTT GAA GAT CAT GTA 120 40 60 GCA CAC AAG AGT GAG GTT GCT CAT CGG TTT AAA GAT TTG GGA GAA GAA AAT TTC AAA GCA ACT CTT A T L GCT A GAT GAG TTA TGC ACA GTT L C T V TTG GTG TTG ATT GCC TTT GCT CAG TAT CTT CAG CAG TGT CCA GCT A CAA GAA 团 AAA ACA TGT GTT K T C V Ø GAG ACA TE T 181 AAT TGT GAC AAA TCA CTT CAT ACC CTT TTT GGA GAC AAA 61 N C D K S L H T L F G D K GAC TGT GCA AAA D C C A K Ω GAA TTT GCA F CAT GAC AAT (H D N I GAA E GAA ATG GCT E M A AAT GAA GTA ACT N E V T ATG TGC ACT GCT TTT M C T A F GAA ACC TAT GGT Ö GTG V 闰 61 GCC 21 A GAT AAA K CGT GAT D Д 121 301 241 361 121 421 141

Figure 15A

540 180 AAA TGT 600 K C 200 CGC CTG AGC 660 R L S 220 240 TGT GAA 840 C E 280 300 GAT CTT ACC AAA 720 GAC AGG GCG GAC CTT 780 D R A D L 260 TAT GCT 960 Y A 320 GAT GAG ATG CCT GCT 900 CTG TTG CCA TGC C TCC AGT AAA CTG AAG GAA S S K L K E GTG GCT TGC AAA CAG AGA ACAAGT CTC CAA AAA TTT GGA GAA AGA GCT TTC AAA GCA TGG GCA S L Q K F G E R A F K A W A GTT TCC AAG TTA GTG GAA AGT AAG GAT GTT E S K D V AAA GCT GCC AAA TGT GCT GAT TTG GAA AAA TCC CAC TGC ATT GCC GAA GTG GAA AAT GAT TCT TCG CAC ACG GAA TGC TGC CAT GGA GAT CTG CTT GAA H T E C C H G D L L E GCT GCT Ø ß TCG ATC TS I S GCTAAG K TGC CAA TTT GCA GAA $_{
m GLL}$ Ø 团 9999 ATC TGT GAA AAT CAG GAT ď $_{\mathrm{TGT}}$ CTT CGG GAT GAA TTA GCT GCT GAT لترا ACA GAA CCC AAA GCT GAG 团 ø GCT GAA TCA AAG TAT P K Y J GCT GAT CTGCCI $_{
m LLL}$ AAA CCTAGA 601 GCC 201 A 901 GAC 301 D AAG K GTC CAG 481 161 541 181 721 781 261 221 661

Figure 15B

AGG CAT CCT GAT 1020 R H P D 340 GAG AAG TGC 1080 E K C 360 GCC GCT GCA GAT CCT CAT GAA TGC TAT GCC AAA GTG TTC GAT GAA TTT AAA CCT CTT 1140 A A A D P H E C Y A K V F D E F K P L 380 1201 TAC AAA TTC CAG AAT GCG CTA TTA GTT CGT TAC ACC AAG AAA GTA CCC CAA GTG TCA ACT 1260 401 Y K F Q N A L L V R Y T K K V P Q V S T 420 1261 CCA ACT CTT GTA GAG GTC TCA AGA AAC CTA GGA AAA GTG GGC AGC AAA TGT TGT AAA CAT 1320 $421~\mathrm{P}$ T L V E V S R N L G K V G S K C C K H 4401321 CCT GAA GCA AAA AGA ATG CCC TGT GCA GAA GAC TAT CTA TCC GTG GTC CTG AAC CAG TTA 1380 441 P E A K R M P C A E D Y L S V V L N O L 460 1381 TGT GTG TTG CAT GAG AAA ACG CCA GTA AGT GAC AGA GTC ACA AAA TGC TGC ACA GAG TCC 1440 GAA GAG CCT CAG AAT TTA ATC AAA CAA AAC TGT GAG CTT TTT GAG CAG CTT GGA GAG 1200 480 GTC GTG CTG CTG AGA CTT GCC AAG ACA TAT GAA ACC ACT CTA V V L L L R L A K T Y E T T L AGA ; TTT TTG TAT GAA TAT GCA a GGC ATG GAG GCA AAG GAT GTC TTC CTG $ext{TCT}$ TGT 1081 ' 1021 341 401 381

Figure 15C

1441 TTG GTG AAC AGG CGA CCA TGC TTT TCA GCT CTG GAA GTC GAT GAA ACA TAC GTT CCC AAA 1500 $481~\mathrm{L}$ V N R R P C F S A L E V D E T Y V P K 500

1501 GAG TTT AAT GCT GAA ACA TTC ACC TTC CAT GCA GAT ATA TGC ACA CTT TCT GAG AAG GAG 1560 Ħ 501 1561 AGA CAA ATC AAG AAA CAA ACT GCA CTT GTT GAG CTT GTG AAA CAC AAG CCC AAG GCA ACA 1620 521 R Q I K K Q T A L V E L V K H K P K A T 540

1621 AAA GAG CAA CTG AAA GCT GTT ATG GAT GAT TTC GCA GCT TTT GTA GAG AAG TGC TGC AAG 1680 260 541

1681 GCT GAC GAT AAG GAG ACC TGC TTT GCC GAG GAG GGT AAA AAA CTT GTT GCT GCA AGT CAA 1740 561 A D D K E T C F A E E G K K L V A A S Q 580

1741 GCT GCC TTA GGC TTA TAA CAT CTA CAT TTA AAA GCA TCT CAG 1782

581 A

Figure 15D